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Key indicators

Single-crystal X-ray study $T = 120 K$ Mean σ (C-C) = 0.003 Å R factor = 0.036 wR factor = 0.093 Data-to-parameter ratio = 9.3

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

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1,2,3,4-Tetra-O-acetyl-a-D-glucopyranuronamide

The title compound, $C_{14}H_{19}NO_{10}$, forms strong hydrogen bonds via amido groups in the [100] direction; soft $C-H\cdots O$ bonds also act to give chains along both [100] and [001], which combine to form a two-dimensional layer.

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Comment

Extensive intermolecular hydrogen-bonding interactions, involving both the hydroxyl and amido groups, are exhibited in the solid-state structure of α -D-glucopyranuronamide, (1) (Flippen & Gilardi, 1974): in the monohydrate $(1) \cdot H_2O$, the water molecule additionally takes part in the strong hydrogen bonding linking the molecules (Perrier & Bryn, 1982). As well as the classic, or hard, hydrogen-bonding interactions of hydroxyl and amido groups, e.g. $N-H\cdots$ O and $O-H\cdots$ O, hydrogen-bonding interactions of a softer nature, such as $C-\frac{1}{2}$ $H \cdots$ O, are found in (1) $\cdot H_2$ O. In general, there is increasing interest in soft hydrogen bonds and their influence on solidstate assemblies (Desiraju, 1991, 1996; Steiner & Desiraju, 1998; Steiner, 1997).

In $1,2,3,4$ -tetra-O-acetyl- α -D-glucopyranuronamide, (2), intermolecular hydrogen bonding falls into two categories: I, strong hydrogen-bonding interactions, $N1-H1B\cdots O6$ ⁱ [symmetry code: (i) $x-1/2$, $-y+5/2$, $-z$], involving just the amido group atoms, and II, a series of soft $C-H\cdots O$ interactions (Table 2). The strong hydrogen bond results in chains in the [100] direction *via* operation of a $2₁$ screw axis (Fig. 2).

The soft $C-H\cdots O$ bonds all involve carbonyl O atoms as acceptors, and either $H-C(pyranose ring)$ or $CH₃H$ atoms as donors, resulting in the formation of an overall two-dimensional sheet (Fig. 3). In this compound, the H atoms in the acetyl group are more acidic than those in simple alkanes due to the electron-withdrawing effects of the adjacent carbonyl group, and hence it is valid to consider the $CH₃$ groups as Hatom donors. Three soft hydrogen bonds, namely $C2 H2\cdots O9^{ii}$, $C8-H8C\cdots O7^{iii}$ and $C14-H14C\cdots O10^{iii}$, form chains via simple translation in the [100] direction (symmetry

The molecular structure of (2), showing 50% probability ellipsoids.

codes are in Table 2). Another soft hydrogen bond, C14 $H14A\cdots$ O6^v also forms chains along [100] *via* operation of a 2₁ screw axis. Finally, a chain forms from C12 $-H12C\cdots$ O7^{iv} interactions in the [001] direction, again *via* action of a $2₁$ screw axis. The intramolecular hydrogen bonds are also listed in Table 2.

Figure 2

Chains formed by the strong hydrogen bond, $N1-H1B\cdots$ O6ⁱ, in the [100] direction. [Symmetry code: (i) $x-1/2$, $-y+5/2$, $-z$.]

Figure 3

A two-dimensional sheet formed by soft hydrogen bonds, $C2-H2\cdots Q9^{ii}$ $CS-H8C\cdots O7^{iii}$ and $C14-H14C\cdots O10^{iii}$, in the [100] and [100] directions, C14 $-H14A\cdots$ O6^v along [100] and C12 $-H12C\cdots$ O7^{iv} along [001]. [Symmetry codes: (ii) $x-1$, y, z; (iii) $x+1$, y, z; (iv) $-x+1/2$, $-y+2$, $z-1/2$; (v) $x+1/2$, $y+5/2$, $-z$.]

The Cremer and Pople puckering parameters, $Q =$ 0.570 (2) \AA , $\theta = 13.6$ (2)^o and $\varphi = 307.0$ (9)^o (Cremer & Pople, 1975), indicate that the conformation of the pyranose ring in (2) is near ${}^{4}C_1$, with a slight distortion in the direction towards E_1 (Koll *et al.*, 1994).

Experimental

Acetic anhydride (10 ml) was added to a solution of p -glucopyranuronamide (1.00 g, 5.2 mmol) in pyridine (10 ml). The reaction mixture was stirred overnight and rotary evaporated. The syrupy residue was crystallized from diethyl ether/petroleum ether (40–60), m.p. 418-421 K, literature value 420-422 K (Barton et al., 1994); yield 1.53 g, 82%. ¹H NMR (250 MHz, CDCl₃): δ 2.05 (s, 6H, 2Me), 2.08 (s, 3H, Me), 2.20 (s, 3H, Me), 4.30 (d, 1H, $J = 10.1$ Hz, H-5), 5.09 (dd, 1H, $J = 3.7, 10.1$ Hz, H-2), 5.25 (t, 1H, $J = 10$ Hz, H-4), 5.54 (t, 1H, $J =$ 10.0 Hz, H-3), 6.26 (br s, 1H, NH)*, 6.38 (d, 1H, $J = 3.7$ Hz, H-1), 6.52 $(br s, 1H, NH)^*$. The two NH signals can appear as a single broad peak in moist CDCl₃. ¹³C NMR (63 MHz, CDCl₃): δ 20.4, 20.6, 20.8, 68.9, 70.1, 70.3, 88.3, 168.9, 169.3, 169.76, 176.79, 169.9. IR (KBr): $(cm⁻¹) 3480, 3230,1748, 1668, 1417, 1383, 1220, 1175, 1134, 1041, 943,$ 563. MS(ES⁺): 384.1 [100%, $M + \text{Na}$]⁺, 745.2 [50%, 2 $M + \text{Na}$]. MS (ES⁻): 360.2 [100%, $M-H$]⁻. X-ray data were collected by the EPSRC National Crystallography Service at the University of Southampton, England, and MS data were obtained by the EPSRC National Mass Spectrometry Service Centre at the University of Wales, Swansea.

Crystal data

Absorption correction: multi-scan (SORTAV; Blessing, 1995, 1997) $T_{\text{min}} = 0.764, T_{\text{max}} = 0.992$

10 467 measured reflections

Refinement

Table 1

$O5-Cl$	1.410(2)	$C4 - O4$	1.435(2)
$O5-C5$	1.432(2)	$C4 - C5$	1.534(3)
$C1 - O1$	1.429(2)	$C5-C6$	1.517(3)
$C1 - C2$	1.530(3)	$C6 - O6$	1.231(3)
$C2 - O2$	1.446(2)	$C6 - N1$	1.332(3)
$C2 - C3$	1.522(3)	$N1 - H1A$	0.92(3)
$C3 - O3$	1.454(2)	$N1 - H1B$	0.87(3)
$C3-C4$	1.519(3)		
$C1 - O5 - C5$	112.31(16)	$C4 - C3 - C2$	111.53(17)
$O5 - C1 - O1$	109.78(15)	$O4 - C4 - C3$	106.92(16)
$O5 - C1 - C2$	110.76(15)	$O4 - C4 - C5$	110.79(15)
$O1 - C1 - C2$	108.65(17)	$C3-C4-C5$	108.70(15)
$O2 - C2 - C3$	106.22(17)	$O5 - C5 - C6$	110.20(17)
$O2 - C2 - C1$	110.57(15)	$O5 - C5 - C4$	107.48(15)
$C3-C2-C1$	113.58(17)	$C6 - C5 - C4$	112.21(16)
$O3 - C3 - C4$	107.74(15)	$N1 - C6 - C5$	116.85(19)
$O3 - C3 - C2$	105.76(16)		

Table 2

Hydrogen-bonding geometry (\mathring{A}, \degree) .

$D - H \cdots A$	$D-H$	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - H \cdots A$
$N1 - H1B \cdots 06$ ¹	0.87(3)	1.99(3)	2.840(3)	165(3)
$C2-H2\cdots O9n$	1.00	2.29	3.168(3)	146
$C8 - H8C \cdots O7$ ⁱⁱⁱ	0.98	2.55	3.366(3)	141
$C12-H12C\cdots O7iv$	0.98	2.60	3.392(3)	138
$C14 - H14A \cdots O6^V$	0.98	2.57	3.529(3)	166
$C14 - H14C \cdots O10$ ⁱⁱⁱ	0.98	2.46	3.283(3)	142
$N1 - H1A \cdots O5$	0.92(3)	2.29(3)	2.658(2)	103(2)
$C3-H3\cdots$ O9	1.00	2.36	2.734(3)	101
$C4 - H4 \cdots O10$	1.00	2.32	2.682(2)	100
$C5 - H5 \cdots O1$	1.00	2.36	2.776(2)	104

Symmetry codes: (i) $x - \frac{1}{2}, \frac{5}{2} - y, -z$; (ii) $x - 1, y, z$; (iii) $1 + x, y, z$; (iv) $\frac{1}{2}$ – x, 2 – y, z – $\frac{1}{2}$; (v) $\frac{1}{2}$ + x, $\frac{5}{2}$ – y, –z.

It was not possible to determine the absolute configuration of the molecule, and thus the conformation of the precursor sugar was adopted. Friedel pairs were averaged. Most H atoms were placed geometrically and refined using a riding model. The exceptions were H atoms in the amido group (N1), which were located from a difference map and refined freely.

Data collection: DENZO (Otwinowski & Minor, 1997) and COLLECT (Hooft, 1998); cell refinement: DENZO and COLLECT; data reduction: DENZO and COLLECT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEX in OSCAIL (McArdle, 1994, 2000) and ORTEPIII for Windows (Farrugia, 1997); software used to prepare material for publication: PLATON (Spek, 2000) and SHELXL97.

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